

Raw Sequence Listing Error Summary

#23
Bill
3/30/99

ERROR DETECTED SUGGESTED CORRECTION

SERIAL NUMBER: 09/093,922B

ATTN: NEW RULES CASES: PLEASE DISREGARD ENGLISH "ALPHA" HEADERS, WHICH WERE INSERTED BY PTO SOFTWARE

- 1 ☒ **Wrapped Nucleics** The number/text at the end of each line "wrapped" down to the next line.
This may occur if your file was retrieved in a word processor after creating it.
Please adjust your right margin to .3, as this will prevent "wrapping".
- 2 ☐ **Wrapped Aminos** The amino acid number/text at the end of each line "wrapped" down to the next line.
This may occur if your file was retrieved in a word processor after creating it.
Please adjust your right margin to .3, as this will prevent "wrapping".
- 3 ☐ **Incorrect Line Length** The rules require that a line not exceed 72 characters in length. This includes spaces.
- 4 ☐ **Misaligned Amino Acid Numbering** The numbering under each 5th amino acid is misaligned. This may be caused by the use of tabs between the numbering. It is recommended to delete any tabs and use spacing between the numbers.
- 5 ☐ **Non-ASCII** This file was not saved in ASCII (DOS) text, as required by the Sequence Rules.
Please ensure your subsequent submission is saved in ASCII text so that it can be processed.
- 6 ☐ **Variable Length** Sequence(s) ____ contain n's or Xaa's which represented more than one residue.
As per the rules, each n or Xaa can only represent a single residue.
Please present the maximum number of each residue having variable length and indicate in the (ix) feature section that some may be missing.
- 7 ☐ **PatentIn ver. 2.0 "bug"** A "bug" in PatentIn version 2.0 has caused the <220>-<223> section to be missing from amino acid sequence(s) _____. Normally, PatentIn would automatically generate this section from the previously coded nucleic acid sequence. Please manually copy the relevant <220>-<223> section to the subsequent amino acid sequence. This applies primarily to the mandatory <220>-<223> sections for Artificial or Unknown sequences.
- 8 ☐ **Skipped Sequences (OLD RULES)** Sequence(s) ____ missing. If intentional, please use the following format for each skipped sequence:
(2) INFORMATION FOR SEQ ID NO:X:
(i) SEQUENCE CHARACTERISTICS: (Do not insert any headings under "SEQUENCE CHARACTERISTICS")
(xi) SEQUENCE DESCRIPTION: SEQ ID NO:X:
This sequence is intentionally skipped

Please also adjust the "(iii) NUMBER OF SEQUENCES:" response to include the skipped sequence(s).
- 9 ☐ **Skipped Sequences (NEW RULES)** Sequence(s) ____ missing. If intentional, please use the following format for each skipped sequence.
<210> sequence id number
<400> sequence id number
000
- 10 ☐ **Use of n's or Xaa's (NEW RULES)** Use of n's and/or Xaa's have been detected in the Sequence Listing.
Use of <220> to <223> is MANDATORY if n's or Xaa's are present.
In <220> to <223> section, please explain location of n or Xaa, and which residue n or Xaa represents.
- 11 ☐ **Use of <213> Organism (NEW RULES)** Sequence(s) ____ are missing this mandatory field or its response.
- 12 ☐ **Use of <220> Feature (NEW RULES)** Sequence(s) ____ are missing the <220> Feature and associated headings.
Use of <220> to <223> is MANDATORY if <213> ORGANISM is "Artificial" or "Unknown"
Please explain source of genetic material in <220> to <223> section.
(See "Federal Register," 6/01/98, Vol. 63, No. 104, pp. 29631-32) (Sec. 1.823 of new Rules)
- 13 ☐ **PatentIn ver. 2.0 "bug"** Please do not use "Copy to Disk" function of PatentIn version 2.0. This causes a corrupted file, resulting in missing mandatory numeric identifiers and responses (as indicated on raw sequence listing).
Instead, please use "File Manager" or any other means to copy file to floppy disk.

RAW SEQUENCE LISTING DATE: 03/23/2001
 PATENT APPLICATION: US/09/093,972B TIME: 12:55:56

Input Set : A:\ES.txt
 Output Set : N:\CRF3\03232001\I093972B.raw

Does Not Comply
 Corrected Diskette Needed

SEQUENCE LISTING

3 (1) GENERAL INFORMATION:
 4 (i) APPLICANT: Nyce, Jonathan W.
 5 (ii) TITLE OF INVENTION: COMPOSITION, FORMULATIONS & METHOD FOR PREVENTION
 6 & TREATMENT OF DISEASES & CONDITIONS ASSOCIATED
 7 WITH BRONCHOCONSTRICTION, ALLERGY(IES) & INFLAMMATION
 E--> 8 (iii) NUMBER OF SEQUENCES: 966 *996 in file (next page)*
 9 (iv) CORRESPONDENCE ADDRESS:
 10 (A) ADDRESSEE: EPIGENESIS PHARMACEUTICALS, INC.
 11 (B) STREET: 7 Clarke Drive
 12 (C) CITY: Cranbury
 13 (D) STATE: New Jersey
 14 (E) COUNTRY: USA
 15 (F) ZIP: 08512
 16 (v) COMPUTER READABLE FORM:
 17 (A) MEDIUM TYPE: Floppy disk
 18 (B) COMPUTER: IBM PC compatible
 19 (C) OPERATING SYSTEM: PC-DOS/MS-DOS
 20 (D) SOFTWARE: PatentIn Release #1.0, Version #1.30
 21 (vi) CURRENT APPLICATION DATA:
 34 (A) APPLICATION NUMBER: US/09/093,972B
 35 (B) FILING DATE: 09-Jun-1998
 36 (C) CLASSIFICATION:
 37 (viii) ATTORNEY/AGENT INFORMATION:
 38 (A) NAME: Amzel, Viviana
 39 (B) REGISTRATION NUMBER: 30,930
 C--> 40 (C) REFERENCE/DOCKET NUMBER: EPI-00672 *see page 4*
 41 (D)
 42 (ix) TELECOMMUNICATION INFORMATION:
 43 (A) TELEPHONE: 609-409-3035
 44 (B) TELEFAX: 413-254-9245
 45 (C) TELEX:

ERRORED SEQUENCES

9891 (2) INFORMATION FOR SEQ ID NO: 984:
 9892 (i) SEQUENCE CHARACTERISTICS:
 9893 (A) LENGTH: 24 base pairs
 9894 (B) TYPE: nucleic acid
 9895 (C) STRANDEDNESS: single
 9896 (D) TOPOLOGY: linear
 9897 (ii) MOLECULE TYPE: DNA (genomic)
 9898 (xi) SEQUENCE DESCRIPTION: SEQ ID NO: 984:
 E--> 9899 5'-CCG CGC CTC CGC CTG CCG CTT CTG

24

delete - do not show prime marks in the sequence itself

(2) INFORMATION FOR SEQ ID NO:996:

(i) SEQUENCE CHARACTERISTICS:

- (A) LENGTH: 14 base pairs
- (B) TYPE: nucleic acid
- (C) STRANDEDNESS: single
- (D) TOPOLOGY: linear

(ii) MOLECULE TYPE: DNA (genomic)

(xi) SEQUENCE DESCRIPTION: SEQ ID NO:996:

TTT CTT CTG TTC CC

14

last sequence in fileformat error - see item 1 on Enov
summary
sheet

(1)

delete at end of file

PRIOR

- (vii) ~~(vi)~~ PRIORITY APPLICATION DATA:
(A) APPLICATION NUMBER: US 08/472,527
(B) FILING DATE: 7-June-1995

- (vii) ~~(vi)~~ PRIORITY APPLICATION DATA:
PRIOR

edit all entries - these are samples

(viii) ATTORNEY/AGENT INFORMATION:

(A) NAME: Amzel, Viviana

(B) REGISTRATION NUMBER: 30,930

(C) REFERENCE/DOCKET NUMBER: EPI-00672

~~(D)~~ delete REFERENCE/DOCKET NUMBER:
=

VERIFICATION SUMMARY

DATE: 03/23/2001

PATENT APPLICATION: US/09/093,972B

TIME: 12:55:59

Input Set : A:\ES.txt

Output Set: N:\CRF3\03232001\I093972B.raw

L:22 M:220 C: Keyword misspelled or invalid format, [(A) APPLICATION NUMBER:]
 L:23 M:220 C: Keyword misspelled or invalid format, [(B) FILING DATE:]
 L:25 M:220 C: Keyword misspelled or invalid format, [(A) APPLICATION NUMBER:]
 L:26 M:220 C: Keyword misspelled or invalid format, [(B) FILING DATE:]
 L:28 M:220 C: Keyword misspelled or invalid format, [(A) APPLICATION NUMBER:]
 L:29 M:220 C: Keyword misspelled or invalid format, [(B) FILING DATE:]
 L:31 M:220 C: Keyword misspelled or invalid format, [(A) APPLICATION NUMBER:]
 L:32 M:220 C: Keyword misspelled or invalid format, [(B) FILING DATE:]
 L:34 M:220 C: Keyword misspelled or invalid format, [(A) APPLICATION NUMBER:]
 L:35 M:220 C: Keyword misspelled or invalid format, [(B) FILING DATE:]
 L:40 M:220 C: Keyword misspelled or invalid format, [(C) REFERENCE/DOCKET NUMBER:]
 L:9899 M:254 E: No. of Bases conflict, Input:24 Counted:24 SEQ:984
 L:9899 M:320 E: (1) Wrong Nucleic Acid Designator, 3
 L:9899 M:204 E: No. of Bases differ, LENGTH:Input:24 Counted:26 SEQ:984
 L:10021 M:336 W: Invalid Amino Acid Number in Coding Region, SEQ ID:996
 L:8 M:203 E: No. of Seq. differs, : Input 966, Counted 996